Edinburgh, UK 03-07-2018





Structure determination of polycrystalline materials using X-rays and electrons

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Electrons as a radiation source



- Accelerating voltage: 100 to 300 keV
- Wavelength: 0.0251 Å @ 200 keV
- Probe electrostatic potential
- Strong interaction (10⁶ stronger than X-rays)
- Require small samples (< 1 μm)
- High vacuum (<10⁻³ mbar)



Electron 'diffractometer'



Single crystal electron diffraction

Limitations

- Dynamical (multiple) scattering
- Beam damage
- Missing wedge
- Goniometer mechanics
- Filling the gaps:
 - Beam tilt (RED)
 - Precession (ADT, pEDT)
 - Continuous rotation (fast EDT, microED, IEDT, CRED)



• More than 200 structures solved (Yun et al., IUCrJ (2015), 2:267)

Fine slicing using beam tilt (RED)

Zhang *et al.*, Z. Krist. (2010), 225:94 Wan *et al.*, J. Appl. Cryst. (2013), 46:1863



Fine slicing using beam tilt (RED)

Zhang *et al.*, Z. Krist. (2010), 225:94 Wan *et al.*, J. Appl. Cryst. (2013), 46:1863

Continuous rotation method (CRED)

Nederlof *et al.*, Acta Cryst. D (2013), 69:1223 Nannenga *et al.*, Nat. Methods (2014), 11:927 Gemmi *et al.*, J. Appl. Cryst. (2015), 48:718 Wang *et al.*, Chem. Commun., (2017), 53:7018





Example: Mordenite



Zeolite Porous aluminosilicate

 $\mathrm{Si}_{48}\mathrm{O}_{96}$

Orthorhombic *Cmcm a* = 18.11 Å *b* = 20.53 Å *c* = 7.528 Å

Mordenite

Rotate: -43.90° to 58.65° @ 0.45°/s (102.55°) Exposure: 0.5 s, oscillation angle: 0.23°





Data collected with M.O. Cichocka (Stockholm University)



100

 $\sqrt{(I_{calc})}$

150

080 8

50

332 061 o

404 006

440 **•**

50

0

Data reduction	XDS
Compl. (<i>Cmcm</i>)	93.6 %
I/σ	6.25
Resolution	0.80 Å
R _{meas}	0.108
R _{obs}	0.088
R _{exp}	0.087
Refinement	ShelXL
Reflections (unique)	1585
Reflections $(F_0 > 4\sigma(F_0))$	1140
$R1 (F_o > 4\sigma(F_o))$	0.158
$R1 (F_o > 4\sigma(F_o))$ R1 (all)	0.158 0.175
$R1 (F_o > 4\sigma(F_o))$ R1 (all) Parameters	0.158 0.175 96
$R1 (F_o > 4\sigma(F_o))$ R1 (all) Parameters Restraints	0.158 0.175 96 0

Structure refinement

Refinement by M.O. Cichocka (Stockholm University)



Structure determination using X-rays and electrons



SSZ-45; S. Smeets et al., Chem. Mater., 2014



SSZ-61; S. Smeets *et al., Angew. Chem.,* 2014



SSZ-87; S. Smeets *et al., J. Am. Chem. Soc.*, 2015





CIT-13; J.H. Kang *et al.*, *Chem. Mater.*, 2017



SCM-14; Y. Luo et al., Chem.-Eur. J., 2017



Outline

- Zeolite IM-18
 - RED + HRTEM + XRPD
- Zeolite SSZ-70
 - HRTEM + XRPD + NMR
- Serial electron diffraction
 - Structure determination
 - Phase analysis
 - Screening







Zeolite IM-18

M.O. Cichocka, Y. Lorgouilloux, S. Smeets, J. Su, W. Wan, P. Caullet, N. Bats, L.B. McCusker, J.-L. Paillaud, and X. Zou. *Cryst. Growth Des.*, 18(4):2441-2451, 2018

Germanosilicate IM-18

Y. Lorgouilloux, et al. French patent 2,923,477 (2007)



4-Dimethylaminopyridine (DMAP)







M. O. Cichocka et al., Cryst. Growth Des., 18(4):2441-2451, 2018

Rotation electron diffraction



Tilt range (°)	119.46 (-66.83 to 52.63)
Tilt step (°)	0.2°
Exposure time/frame (s)	1.0
No. of frames	649
Crystal size (μm)	0.66 x 0.74
Resolution (Å)	1.05
Completeness (%)	89.9
Reflections	1265

Index Bragg spots

Orthorhombic <i>l</i>	'mma / Im2a
a = 531 Å	$\alpha = 89.79^{\circ}$

u – 3.31 A	u = 09.79
<i>b</i> = 15.07 Å	<i>b</i> = 88.81°
<i>c</i> = 17.06 Å	γ = 90.35°

hkl: h + k + l = 2nhk0: h = 2n, k = 2n

M. O. Cichocka et al., Cryst. Growth Des., 18(4):2441-2451, 2018

Average framework structure from SHELXS



Average framework structure from SHELXS



Monoclinic *P*2₁/*m a* = 10.336 Å, *b* = 14.984 Å, *c* = 17.734 Å, *β* = 106.94°

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Selected area electron diffraction





M. O. Cichocka et al., Cryst. Growth Des., 18(4):2441-2451, 2018

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HRTEM



M. O. Cichocka et al., Cryst. Growth Des., 18(4):2441-2451, 2018

HRTEM



M. O. Cichocka et al., Cryst. Growth Des., 18(4):2441-2451, 2018



M. O. Cichocka et al., Cryst. Growth Des., 18(4):2441-2451, 2018

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Summary IM-18

- Structure of IM-18 determined by combining methods
 - RED \rightarrow Average structure
 - $\text{ SAED } \rightarrow \text{Disorder}$
 - HRTEM \rightarrow Short-range order
 - XRPD \rightarrow Structure completion
 - \rightarrow Model validation
- New zeolite framework topology
- Experimental evidence for 2D stacking disorder



Zeolite SSZ-70

S. Smeets, Z.J. Berkson, D. Xie, S.I. Zones, W. Wan, X. Zou, M.-F. Hsieh, B.F. Chmelka, L.B. McCusker, and C. Baerlocher. *J. Am. Chem. Soc.*, 139(46):16803-16812, 2017

Zeolite SSZ-70

Stacey Zones and Alan Burton, US Patent 7,108,843 B2 (2006) Molecular sieve SSZ-70 composition of matter and synthesis thereof







Runnebaum et al., 2014, ACS Catal., 4, 2364

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Electron diffraction (as-made)



Along [001]

Along [100]



HRTEM (as-made)

Stacking disorder along [001]



MWW-layers



Stacking faults



Collected by Wei Wan, Stockholm University, SE

Solid-state ²⁹Si MAS NMR



Hsieh, Aronson and Chmelka (2014)







Archer et al., **2010**, *Micropor*. *Mesopor*. *Mat.*, 130, 255 Camblor et al., **1998**, *J. Phys. Chem. B*, 102, 44

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Solid-state ²⁹Si MAS NMR



Hsieh, Aronson and Chmelka (2014)

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Disorder model





Random arrangement of **MWW** layers

 $P(A \rightarrow A) = 0\%$ $P(A \rightarrow B) = 50\%$ $P(A \rightarrow C) = 50\%$







Simulations using DiFFaX

DiFFaX: Traecy et al., 1991, Proc. R. Soc. Lond. A, 433, 499

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Interlayer region











Zach Berkson and Brad Chmelka, UC Santa Barbara, USA

ppm







Zach Berkson and Brad Chmelka, UC Santa Barbara, USA

Interlayer region





Model 2

50%

Structure of calcined SSZ-70



Summary SSZ-70

- Structure of SSZ-70 determined by combining methods
 - − HRTEM \rightarrow Short-range order
 - XRPD \rightarrow Long-range order
 - − 2D NMR \rightarrow Nanostructure
- New stacking arrangement of **MWW**-layers
- Mixed silanol sites at the nanoscale can help explain enhanced catalytic behaviour of SSZ-70



Serial electron diffraction

Serial electron diffraction





Randomly oriented crystals 1 crystal = 1 diffraction pattern Combine data from many crystals

Collect data ~3000 crystals/hour











GUI for data collection

ps: 13.57 interval (ms): 73.68 🗌 Increase size 🖂 A	o contrast						
exposure (s) 0.05 🜩 Brightness 1.0 🜩 DisplayRange 11800	Directory:	C:\insta	amatic\wor	rk_2018-0	6-13		Browse.
	Sample name:	experim	nent				1
	Flatfield:	C:\insta	amatic\flatf	field.tiff			Browse.
	Open wor	k directo	ory	Open se	ettings directory	Delete last	experiment
	CRED autoCRED) serialE otation e ne: [D RED ctr electron dif	rl learnir fraction	ng expert about	🗌 Beam ur	nblanker
	Image interv	ral:	10	÷		Enable in	mage interv
	Diff derocus	;	0.01			Relay	heam
	Select outpu	it format	ts:		☑ .tiff ☑ DIALS (.smv)	☑ XDS (.sr ☑ REDp (.	mv) mrc)
	Start Collection		s	Stop Collection			
Save image							

Data collection (Zeolite Y)

images\image_0000.h5



data\image_0000_0000.h5



Data collection (zeolite A)



Diffraction

Collect data

2 µm

Total: 1107 patterns

Serial electron diffraction

- Structure determination?
 - Phase analysis?
 - Screening?

Structure determination: orientation finding

- Forward projection model using known lattice parameters
- Generate pattern library of all possible orientations (~1.5M in P1)
- Match best orientation and index data



Source code: www.github.com/stefsmeets/problematic

Based on: Rius et al., IUCrJ (2015), 2:452

Structure determination: Data Merging

Challenges

- Scaling
- Dynamical effects
- Reflection partiality

SerialMerge – rank-based merging

- Avoid scaling
- Avoid modelling intensities
- Robust with low quality data



S. Smeets & W. Wan, *J. Appl. Cryst.* (2017). **50**, 885-892 www.github.com/stefsmeets/serialmerge

Structure determination



Zeolite A $Fm\overline{3}c$ a = 24.61 Å $Si_{96}Al_{96}O_{384}$ *Z* = 192

200 frames

OK SIMU Reflections Total: 19804 Unique: 227 d_{min}: 1.03 Å Compl.: 100%

Structures solved



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Serial electron diffraction

- Structure determination?
- ➡ Phase analysis?
 - Screening?

Phase analysis: Co-CAU-36

images\image_0342.h5

data\image_0342_0002.h5

Scan 200 x 200 µm in 30 minutes 1202 diffraction patterns

Sample from Bin Wang & Ken Inge (Stockholm University)

Phase analysis: Co-CAU-36



data\image_0280_0012.h5



data\image_0468_0003.h5





data\image_0450_0002.h5



data\image_0469_0001.h5









1202 diffraction patterns500 contained reflections -> 6 impurity crystals

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Phase analysis: Co-CAU-36





34.45 to -13.79°Oscillation angle: 0.23°1.5 min data collection



XDS

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'Quantitative' phase analysis



Co-CAU-36: ~99% 494 patterns CoO (wurtzite): ~1% 6 patterns

Serial electron diffraction

- Structure determination?
- Phase analysis?
- ➡ Screening?

Screening: Mordenite

images\image_0074.h5





Scan 200 x 200 μm in 24 minutes 836 diffraction patterns (2090 / hour)



Screening: Crystal selection

Crystal selection

- 1. Find isolated crystals
 - Must be 0.5 µm away from edge
 - No crystals in 1.5 μ m radius
- 2. Select most suitable crystals
 - Machine learning (CNN)

Screening: Machine learning

• A deep convoluted neural network trained on ~78.000 diffraction patterns predicts which crystals are suitable for collecting CRED data



Prediction: 1.0





Prediction: 1.0





Prediction: 0.26





Prediction: 0.25



Jonas Ångström (Stockholm University)

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Screening: 6 of the 'best' crystals (53)



frame: 230, crystal: 1, size: 0.060 µm²



frame: 104, crystal: 1, size: 0.163 μm^2



frame: 252, crystal: 1, size: 0.350 μm^2



frame: 188, crystal: 2, size: 0.351 μm^2



frame: 419, crystal: 1, size: 0.441 μm^2



Screening: 6 of the 'worst' crystals (53)



frame: 400, crystal: 2, size: 0.069 µm²





frame: 430, crystal: 1, size: 0.121 μm^2



Prediction: 0.0

frame: 392, crystal: 2, size: 0.250 µm²

frame: 449, crystal: 3, size: 0.040 μm^2





Automated data collection



Rotation: -44.0 to 47.4° @ 0.76°/s (91.4°) Exposure: 0.5 s, oscillation angle: 0.39°



Data collected by Bin Wang (Stockholm University)

Conclusions

- Electrons are very well suited for structure determination
 - Reliable crystal structures can be obtained
- PXRD data are valuable for
 - Structure validation against bulk material
 - Structure completion (*e.g.* cations/templates/adsorbants in zeolites)
- Combination of methods is essential to find all the details
- SerialED data can be collected routinely & automatically
 - Structure determination
 - Phase analysis
 - Screening
- Future: Combined SerialED and CRED for automated crystal picking and data collection